AMENDMENTS TO SPECIFICATION UNDER THE PROVISION OF 37 CFR § 1.121(b)(2)(i)

Please insert the following section on page 1, immediately below the title:

Cross Reference to Related Application

This application claims the benefit of U.S. Provisional Application No. 60/397,758, filed 23 July 2002.

Please delete the paragraph starting at page 13, line 20 and insert the following therefor:

A preferred method of providing a T cell epitope, therefore, involves computational methods of identifying MHC class I or class II binding peptides, such as computational "threading" algorithms (e.g., see Altuvia et al., 1995). For example, computer analysis may be conducted using MPT (ver 1.0) software (Biovation, Aberdeen, UK), which performs peptide threading according to the methods disclosed by Fothergill et al., 1998, wherein, an index of potential peptide binding to 18 different MHC class II DR alleles (covering greater than 96% of the HLA-DR allotypes extant in the human population) may be calculated. Alternatively, or in concert, a comparison of suspected epitope sequences may be made against preexisting databases of MHC-binding motifs (e.g., "MHCPEP: A database of MHC binding peptides, v.1.3", Brusic, 1998, http://wehih.wehi.edu.au/mhcpep/).